

<u>Set Name</u>	<u>Query</u>	<u>Hit Count</u>	<u>Set Name</u>
side by side			result set
<i>DB=DWPI; PLUR=YES; OP=ADJ</i>			
<u>L19</u>	L18 and l15 and l16	0	<u>L19</u>
<u>L18</u>	L17 near10 l14	17	<u>L18</u>
<u>L17</u>	recombin\$ or variant	53688	<u>L17</u>
<u>L16</u>	library	12726	<u>L16</u>
<u>L15</u>	heteroduplex	168	<u>L15</u>
<u>L14</u>	mismatch adj repair	83	<u>L14</u>
<i>DB=PGPB; PLUR=YES; OP=ADJ</i>			
<u>L13</u>	L12 and l9 and l10	12	<u>L13</u>
<u>L12</u>	L11 near10 l8	38	<u>L12</u>
<u>L11</u>	recombin\$ or variant	49333	<u>L11</u>
<u>L10</u>	library	27520	<u>L10</u>
<u>L9</u>	heteroduplex	1652	<u>L9</u>
<u>L8</u>	mismatch adj repair	912	<u>L8</u>
<i>DB=USPT; PLUR=YES; OP=ADJ</i>			
<u>L7</u>	l6 and l3 and l2	26	<u>L7</u>
<u>L6</u>	l1 near10 l5	84	<u>L6</u>
<u>L5</u>	L4 or variant	170307	<u>L5</u>
<u>L4</u>	recombin\$	75498	<u>L4</u>
<u>L3</u>	library	53319	<u>L3</u>
<u>L2</u>	heteroduplex	2284	<u>L2</u>
<u>L1</u>	mismatch adj repair	576	<u>L1</u>

END OF SEARCH HISTORY

100 10.03 01:45P

INFORMATION SPECIALTIES 793 931 4591
Novozymes North America

P.02

#1
3/17/03
RM

Attorney Docket No: 10070.200-US

PATENT

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of Jesper Vind

Confirmation No: 9234

Serial No 10/004,963

Group Art Unit 1645

Filed December 4, 2001

Examiner: Ponnaluri, P.

Re: Method For Producing a Polynucleotide Library

POWER TO INSPECT

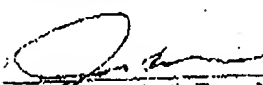
Assistant Commissioner for Patents
Washington, DC 20231

S:

Please permit Information Specialties Corp., Gary Wasserman or their representatives, to inspect the above-captioned application and to make copies of any of the papers that they may desire.

Respectfully submitted,

Date: March 13, 2002


Jason I. Garbell, Reg. No. 44,116
Novozymes North America, Inc.
500 Fifth Avenue, Suite 1600
New York, NY 10110
(212) 840-0097

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Does Not Comply
Corrected Diskette Needed

10/004993

~~ CProject

CProjectData -Method for Producing a Polynucleotide Library

10070.200-US

____-____-____ %S:\ENZYME\PAT-ADM\PWL\10070\Sequences ~~

CDNASequence Seq1 Artificial sequenceIgaatgacttg gttgacgcgt caccagtcac
30 agctrymkswbdhvn DNA ~~ CCommentFeature
Primer ...`a 0aa ~~ COtherFeature

misc_feature 5' Phosphorylation 1 1 ? Seq2 Artificial
sequence cttattagtaggttggtacttcgag agctrymkswbdhvn DNA
? Primer " ^ ' o ^ ' ?